Please amend the application as follows:

In the specification:

Replace Table 1 on page 15 with the amended Table below.

Table 1. Nucleotide sequence of primers used to generate promoter fragments

Primer No.	Restriction site	Promoter sequence	Position	Sequence 5'-3'	
1582	HindIII (AAG CTT)	SEQ ID NO: 1	3440-3424	AAG CTT CTC GGC GCG CGG GCC CC	.
				(SEQ ID NO: 3)	
1583	NheI (GCT AGC)	SEQ ID NO: 1	2341-2362	GCT AGC CAA GAG CTT CTG GAG CC	ēG ∵
	•.			(SEQ ID NO: 4)	**************************************
1584	Nhel (GCT AGC)	SEQ ID NO: 1	720-741	GCT AGC TGT TAC ATG CAG AGC AA	AT .
-				C (SEQ ID NO: 5)	
1585	HindIII (AAG CTT)	SEQ ID NO: 2	4439-4421	AAG CTT CCT ACG GCC CCC GCG	
				(SEQ ID NO: 6)	**
1586	Nhel (GCT AGC)	SEQ ID NO: 2	3321-3340	GCT AGC GCG CAC TGC AAT GCC CT	.c
				(SEQ ID NO: 7)	

Replace Table 2 on pages 20 and 21 with the amended Table below.

Table 2. Oligonucleotide primer used in the mutagenesis experiments

Primer		And the second s
ET TIMET		Primer Sequences
	41 Te	
1949 P R1b Cre Fwd		CGCCGCCGT TTC GTCAGAGCCCCCT
1343 I KID CIE IWG		COCCOCCCITICAGAGCCCCCI
		(SEQ ID NO: 8)
		ie:
1950 P R1b Cre Rev		AGGGGGCTCTGAC CAA ACGGGCGGCG
1930 F KID CIE REV		
·		
		(SEQ ID NO: 9)
1951 P Rla GCI Fwd		CTCTCTTCCCCCCTAACTGCCTTCCC
		ISEO ID NO. 10)
		(SEQ ID NO: 10)
1952 P Rla GCI Rev		GGGAAGGCAG TTA GGGGGGAAGAGAG
1304 1 1124 002 110		
		(SEQ ID NO: 11)
		1000 100 100
1052 5 51 0017 5 1		COOCOMOCA CHIEF COCCOMOCCA MCC
1953 P Rla GCII Fwd		GGCGGTCCAG TTA GGGGCTGGGATCC
	•••	
		(SEQ ID NO: 12)
		GGATCCCAGCCCC TAA CTGGACCGCC
1954 P Rla GCII Rev		
		(SEQ ID NO: 13)
2051 P Rla GCIII Fwd		CCTCTCCACCGCCCTAACCACCGCGCTGTG
2001 I RIA GOLLI PWQ	Ă.	
	7 1	(GEO TO NO. 14)
· ·		(SEQ ID NO: 14)
		CACAGCGCGGTGGTTAGGGCGGTGGAGAGG
2052 P Rla GCIII Rev		
	- 27	(SEQ ID NO: 15)

2053 P R1b GCIVs Fwd		CCCCAGCTCCCGCCCTAACCCCCCACCCC
		(SEQ ID NO: 16)
	•	*
2054 P R1b GCIVs Rev	***	GGGGTGGGGGTTAGGGCGGGAGCTGGGG
Least I Mile Octive Kev	<i>x</i>	
		(SEQ ID NO: 17)
2055 8 811 887	<u> </u>	
2055 P R1b GCV Fwd		CGCTTCCCTCCCCTAACCCTTCCTGCC
	,	(SEQ ID NO: 18)
2056 P R1b GCV Rev		GGCAGGAAGGG TTA GGGGAGGGAAGCG
		(SEQ ID NO: 19)
2057 P R1b GCVI Fwd		CCCTCCCCTAACCTCCGACTGT
·		(SEQ ID NO: 20)
2058 P R1b GCVI Rev		ACAGTCGGAGG TTA GGGGAGGGGAGGG
		(SEQ ID NO: 21)
2059 P R1b GCVII Fwd		CTCCGCCCACCCC TAA CTCCTGGCAC
		(SEO ID NO: 22)
2060 P R1b GCVII Rev		GTGCCAGGAG TTA GGGGTGGGCGGAG
TOO I MID GOVIE WEA	*	76.4
		(SEQ ID NO: 23)
2146 P R1b GCIVd Fwd		CCCCAGCTCCCTAACTAACCCCCACCCC
		(SEQ ID NO: 24)
2147 p p1h corul p		GGGGTGGGGGTTAGTTAGGGAGCTGGGG
2147 P R1b GCIVd Rev	. 25	
		(SEQ ID NO: 25)
		Tong 10 10 - 20/

Replace the legend to Figure 6 running from page 25, line 24 through page 26, line 3 with the amended legend below.

Figure 6. Identification of nuclear factors binding to the Plb consensus CRE site using CREB/ATF super-shift antibodies.

Nuclear extracts (5µg) from ND7/23 cells were incubated with double-stranded ³²P-labeled oligonucleotides containing the Plb consensus CRE site (sense:5'-CGCCGCCGTGACGTCAGAGCCCCCT-3'(SEQ ID NO: 26)). In lane 1, no antibody was added. In lane 2, a mouse monoclonal antibody (sc-270 Santa Cruz Biotechnology, Santa Cruz, CA) reactive with members of the ATF/CREB family such as ATF-1 p35, CREB-1 p43 and CREM-1 was pre-incubated at room temperature for 20 min before addition of ³²P-labeled probe. The specific complex between nuclear factors and the CRE is indicated by a star and the super-shifted complex is indicated by two stars.

Replace the present Sequence Listing with the revised one on 11 substitute sheets enclosed herewith.